OIPE JC22
OCT 18 2004
PATENT & TRADEMARK OFFICE

Sequence Listing

<110> CHUNTHARAPAI, ANAN
GREWAL, IQBAL
KIM, KYUNG JIN
YAN, MINHONG

<120> TACI Antibodies and Uses Thereof

<130> P1942R1

<140> US 10/626,914
<141> 2003-07-25

<150> US 60/398,530
<151> 2002-07-25

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Pro	Gly	Leu	Lys	Leu	Ser	Ala	Asp	Gln	Val	Ala	Leu	Val	Tyr	Ser	
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185 190 195

Gln Pro Arg Ser Arg Pro Arg Gln Ser Pro Ala Lys Ser Ser Gln
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Asp His Ala Met Glu Ala Gly Ser Pro Val Ser Thr Ser Pro Glu
215 220 225

Pro Val Glu Thr Cys Ser Phe Cys Phe Pro Glu Cys Arg Ala Pro
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Thr Gln Glu Ser Ala Val Thr Pro Gly Thr Pro Asp Pro Thr Cys
245 250 255

Ala Gly Arg Trp Gly Cys His Thr Arg Thr Thr Val Leu Gln Pro
260 265 270

Cys Pro His Ile Pro Asp Ser Gly Leu Gly Ile Val Cys Val Pro
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<213> Homo sapien

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aaagagatct gacacatcat ctattctgac aaaagatttt taaagtggca 200

ctgctcgagt cgaaatggtt aattacctag cagaaattga tttctctatc 250

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<212> PRT
<213> Homo sapien

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Asn Thr Pro Pro Leu Thr Cys Gln Arg Tyr Cys Asn Ala Ser Val
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Thr Asn Ser Val Lys Gly Thr Asn Ala Ile Leu Trp Thr Cys Leu
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Gly Leu Ser Leu Ile Ile Ser Leu Ala Val Phe Val Leu Met Phe
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Leu Leu Arg Lys Ile Ser Ser Glu Pro Leu Lys Asp Glu Phe Lys
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Asn Thr Gly Ser Gly Leu Leu Gly Met Ala Asn Ile Asp Leu Glu
95 100 105

Lys Ser Arg Thr Gly Asp Glu Ile Ile Leu Pro Arg Gly Leu Glu
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Tyr Thr Val Glu Glu Cys Thr Cys Glu Asp Cys Ile Lys Ser Lys
125 130 135

Pro Lys Val Asp Ser Asp His Cys Phe Pro Leu Pro Ala Met Glu
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Glu Gly Ala Thr Ile Leu Val Thr Thr Lys Thr Asn Asp Tyr Cys
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<212> PRT
<213> Homo sapien

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Leu Ala Ser Leu Arg Ala Glu Leu Gln Gly His His Ala Glu Lys
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Leu Pro Ala Gly Ala Gly Ala Pro Lys Ala Gly Leu Glu Glu Ala
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Pro Ala Val Thr Ala Gly Leu Lys Ile Phe Glu Pro Pro Ala Pro
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Gly Glu Gly Asn Ser Ser Gln Asn Ser Arg Asn Lys Arg Ala Val
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Gln Gly Pro Glu Glu Thr Val Thr Gln Asp Cys Leu Gln Leu Ile
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Ala Asp Ser Glu Thr Pro Thr Ile Gln Lys Gly Ser Tyr Thr Phe
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Val Pro Trp Leu Leu Ser Phe Lys Arg Gly Ser Ala Leu Glu Glu
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Lys Glu Asn Lys Ile Leu Val Lys Glu Thr Gly Tyr Phe Phe Ile
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Tyr Gly Gln Val Leu Tyr Thr Asp Lys Thr Tyr Ala Met Gly His
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Leu Ile Gln Arg Lys Lys Val His Val Phe Gly Asp Glu Leu Ser
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Leu Val Thr Leu Phe Arg Cys Ile Gln Asn Met Pro Glu Thr Leu
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Pro Asn Asn Ser Cys Tyr Ser Ala Gly Ile Ala Lys Leu Glu Glu
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Gly Asp Glu Leu Gln Leu Ala Ile Pro Arg Glu Asn Ala Gln Ile
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<213> Homo sapien

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<212> PRT
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Met Ala Leu Leu Thr Gln Gln Thr Glu Leu Gln Ser Leu Arg Arg
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Val Leu Thr Gln Lys Gln Lys Lys Gln His Ser Val Leu His Leu
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Val Pro Ile Asn Ala Thr Ser Lys Asp Asp Ser Asp Val Thr Glu
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Val Met Trp Gln Pro Ala Leu Arg Arg Gly Arg Gly Leu Gln Ala
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Gln Gly Tyr Gly Val Arg Ile Gln Asp Ala Gly Val Tyr Leu Leu
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Tyr Ser Gln Val Leu Phe Gln Asp Val Thr Phe Thr Met Gly Gln
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Val Val Ser Arg Glu Gly Gln Gly Arg Gln Glu Thr Leu Phe Arg
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Cys Ile Arg Ser Met Pro Ser His Pro Asp Arg Ala Tyr Asn Ser
               200                 205                 210

Cys Tyr Ser Ala Gly Val Phe His Leu His Gln Gly Asp Ile Leu
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Ser Val Ile Ile Pro Arg Ala Arg Ala Lys Leu Asn Leu Ser Pro
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His Gly Thr Phe Leu Gly Phe Val Lys Leu
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<212> DNA
<213> Homo sapien

<400> 13

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gccgtgtgga ccaggaggag cgctggtcac tcagctgccg caaggagcaa 100

ggcaagttct atgaccatct cctgagggac tgcatcagct gtgcctccat 150

ctgtggacag caccctaagc aatgtgcata cttctgtgag aacaagctca 200

ggagcccagt gaaccttcca ccagagctca ggagacagcg gagtggagaa 250

gttgaaaaca attcagacaa ctcgggaagg taccaaggat tggagcacag 300

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aggtggccct ggtctacagc acgctggggc tctgcctgtg tgccgtcctc 400

tgctgcttcc tggtggcggt ggcctgcttc ctcaagaaga ggggggatcc 450
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ctgctcctgc cagccccgct caaggccccg tcaaagtccg gccaagtctt 500

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ggagagcgca gtcacgcctg ggacccccga ccccacttgt gctggaaggt 650

gggggtgcca caccaggacc acagtcctgc agccttgccc acacatccca 700

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tgcataaatg ggggtcaggg agggaaagga ggagggagag agatggagag 800

gaggggagag agaaagagag gtggggagag gggagagaga tatgaggaga 850

gagagacaga ggaggcagaa agggagagaa acagaggaga cagagaggga 900

gagagagaca gagggagaga gagacagagg ggaagagagg cagagaggga 950

aagaggcaga gaaggaaaga gacaggcaga gaaggagaga ggcagagagg 1000

gagagaggca gagagggaga gaggcagaga gacagagagg gagagaggga 1050

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<210> 14
<211> 246
<212> PRT
<213> Homo sapien

<400> 14

Met	Ser	Gly	Leu	Gly	Arg	Ser	Arg	Arg	Gly	Gly	Arg	Ser	Arg	Val	
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Asp	Gln	Glu	Glu	Arg	Trp	Ser	Leu	Ser	Cys	Arg	Lys	Glu	Gln	Gly	
				20				25						30	
Lys	Phe	Tyr	Asp	His	Leu	Leu	Arg	Asp	Cys	Ile	Ser	Cys	Ala	Ser	
				35				40						45	
Ile	Cys	Gly	Gln	His	Pro	Lys	Gln	Cys	Ala	Tyr	Phe	Cys	Glu	Asn	
				50				55						60	
Lys	Leu	Arg	Ser	Pro	Val	Asn	Leu	Pro	Pro	Glu	Leu	Arg	Arg	Gln	
				65				70						75	
Arg	Ser	Gly	Glu	Val	Glu	Asn	Asn	Ser	Asp	Asn	Ser	Gly	Arg	Tyr	
				80				85						90	
Gln	Gly	Leu	Glu	His	Arg	Gly	Ser	Glu	Ala	Ser	Pro	Ala	Leu	Pro	
				95				100						105	
Gly	Leu	Lys	Leu	Ser	Ala	Asp	Gln	Val	Ala	Leu	Val	Tyr	Ser	Thr	
				110				115						120	

Leu Gly Leu Cys Leu Cys Ala Val Leu Cys Cys Phe Leu Val Ala
125 130 135

Val Ala Cys Phe Leu Lys Lys Arg Gly Asp Pro Cys Ser Cys Gln
140 145 150

Pro Arg Ser Arg Pro Arg Gln Ser Pro Ala Lys Ser Ser Gln Asp
155 160 165

His Ala Met Glu Ala Gly Ser Pro Val Ser Thr Ser Pro Glu Pro
170 175 180

Val Glu Thr Cys Ser Phe Cys Phe Pro Glu Cys Arg Ala Pro Thr
185 190 195

Gln Glu Ser Ala Val Thr Pro Gly Thr Pro Asp Pro Thr Cys Ala
200 205 210

Gly Arg Trp Gly Cys His Thr Arg Thr Thr Val Leu Gln Pro Cys
215 220 225

Pro His Ile Pro Asp Ser Gly Leu Gly Ile Val Cys Val Pro Ala
230 235 240

Gln Glu Gly Gly Pro Gly
245

<210> 15
<211> 595
<212> DNA
<213> Homo sapien

<400> 15
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cagcccccac gccctgcgtc ccggccgagt gcttcgacct gctggtccgc 100

cactgcgtgg cctgcgggct cctgcgcacg ccgcggccga aaccggccgg 150

ggccagcagc cctgcgccca ggacggcgct gcagccgcag gagtcggtgg 200

gcgcgggggc cggcgaggcg gcgctgcccc tgcccgggct gctctttggc 250

gcccccgcgc tgctgggcct ggcactggtc ctggcgctgg tcctggtggg 300

tctggtgagc tggaggcggc gacagcggcg gcttcgcggc gcgtcctccg 350

cagaggcccc cgacggagac aaggacgccc cagagcccct ggacaaggtc 400

atcattctgt ctccgggaat ctctgatgcc acagctcctg cctggcctcc 450

tcctggggaa gacccaggaa ccaccccacc tggccacagt gtccctgtgc 500

cagccacaga gctgggctcc actgaactgg tgaccaccaa gacggccggc 550

cctgagcaac aatagcaggg agccggcagg aggtggcccc tgccc 595

<210> 16
<211> 184
<212> PRT
<213> Homo sapien

<400> 16

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Met Arg Arg Gly Pro Arg Ser Leu Arg Gly Arg Asp Ala Pro Ala
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Pro Thr Pro Cys Val Pro Ala Glu Cys Phe Asp Leu Leu Val Arg
                20                  25                  30

His Cys Val Ala Cys Gly Leu Leu Arg Thr Pro Arg Pro Lys Pro
                35                  40                  45

Ala Gly Ala Ser Ser Pro Ala Pro Arg Thr Ala Leu Gln Pro Gln
                50                  55                  60

Glu Ser Val Gly Ala Gly Ala Gly Glu Ala Ala Leu Pro Leu Pro
                65                  70                  75

Gly Leu Leu Phe Gly Ala Pro Ala Leu Leu Gly Leu Ala Leu Val
                80                  85                  90

Leu Ala Leu Val Leu Val Gly Leu Val Ser Trp Arg Arg Arg Gln
                95                 100                 105

Arg Arg Leu Arg Gly Ala Ser Ser Ala Glu Ala Pro Asp Gly Asp
               110                 115                 120

Lys Asp Ala Pro Glu Pro Leu Asp Lys Val Ile Ile Leu Ser Pro
               125                 130                 135

Gly Ile Ser Asp Ala Thr Ala Pro Ala Trp Pro Pro Pro Gly Glu
               140                 145                 150

Asp Pro Gly Thr Thr Pro Pro Gly His Ser Val Pro Val Pro Ala
               155                 160                 165

Thr Glu Leu Gly Ser Thr Glu Leu Val Thr Thr Lys Thr Ala Gly
               170                 175                 180

Pro Glu Gln Gln
```

<210> 17
<211> 265
<212> PRT
<213> Homo sapien

<400> 17

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Met Ser Gly Leu Gly Arg Ser Arg Arg Gly Gly Arg Ser Arg Val
  1               5                  10                  15

Asp Gln Glu Glu Arg Phe Pro Gln Gly Leu Trp Thr Gly Val Ala
                20                  25                  30

Met Arg Ser Cys Pro Glu Glu Gln Tyr Trp Asp Pro Leu Leu Gly
                35                  40                  45

Thr Cys Met Ser Cys Lys Thr Ile Cys Asn His Gln Ser Gln Arg
                50                  55                  60

Thr Cys Ala Ala Phe Cys Arg Ser Leu Ser Cys Arg Lys Glu Gln
                65                  70                  75

Gly Lys Phe Tyr Asp His Leu Leu Arg Asp Cys Ile Ser Cys Ala
                80                  85                  90
```

```
Ser Ile Cys Gly Gln His Pro Lys Gln Cys Ala Tyr Phe Cys Glu
                 95                  100                 105

Asn Lys Leu Arg Ser Pro Val Asn Leu Pro Pro Glu Leu Arg Arg
                110                 115                 120

Gln Arg Ser Gly Glu Val Glu Asn Asn Ser Asp Asn Ser Gly Arg
                125                 130                 135

Tyr Gln Gly Leu Glu His Arg Gly Ser Glu Ala Ser Pro Ala Leu
                140                 145                 150

Pro Gly Leu Lys Leu Ser Ala Asp Gln Val Ala Leu Val Tyr Ser
                155                 160                 165

Thr Leu Gly Leu Cys Leu Cys Ala Val Leu Cys Cys Phe Leu Val
                170                 175                 180

Ala Val Ala Cys Phe Leu Lys Lys Arg Gly Asp Pro Cys Ser Cys
                185                 190                 195

Gln Pro Arg Ser Arg Pro Arg Gln Ser Pro Ala Lys Ser Ser Gln
                200                 205                 210

Asp His Ala Met Glu Ala Gly Ser Pro Val Ser Thr Ser Pro Glu
                215                 220                 225

Pro Val Glu Thr Cys Ser Phe Cys Phe Pro Glu Cys Arg Ala Pro
                230                 235                 240

Thr Gln Glu Ser Ala Val Thr Pro Gly Thr Pro Asp Pro Thr Cys
                245                 250                 255

Ala Gly Arg Thr Ala Pro Pro Arg Glu Gly
                260                 265
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